

LN
 RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/826,791

DATE: 04/20/2001
 TIME: 11:49:11

Input Set : A:\Pc10914a.app
 Output Set: N:\CRF3\04202001\I826791.raw

3 <110> APPLICANT: Pfizer Inc
 5 <120> TITLE OF INVENTION: Novel Polypeptide
 7 <130> FILE REFERENCE: PC10914ADAM
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/826,791
 C--> 10 <141> CURRENT FILING DATE: 2001-04-05
 12 <150> PRIOR APPLICATION NUMBER: 0008504.3
 13 <151> PRIOR FILING DATE: 2000-04-05
 15 <150> PRIOR APPLICATION NUMBER: 60/198,367
 16 <151> PRIOR FILING DATE: 2000-04-19
 18 <160> NUMBER OF SEQ ID NOS: 6
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 23 <211> LENGTH: 993
 24 <212> TYPE: DNA
 25 <213> ORGANISM: Homo sapiens
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 32 gggTTGTCCA tATATGTTT CCTGCAGCCT tATAAGAAGT ccACATCTGT gAACGTTTTC 180
 34 atGCTAAATC tGGCCATTTC agATCTCTG ttCATAAAGCA CGCTTCCCTT cAGGGCTGAC 240
 36 tATTATCTTA gAGGCTCCAA ttGGATATTt ggAGACCTGG CCTGCAGGAT tATGCTTAT 300
 38 tcCTGTATG tCAACATGTA cAGCAGTATT tATTCCTGA CGTGTCTGAG tGTTGTGCGT 360
 40 tTCCTGGCAA tGGTTCACCC CTTTCCGGCTT CTGCTATGTC ccAGCATCAG gAGTGCCTGG 420
 42 atCCCTCTGTG gGATCATATG gATCCTTATC atGGCTTCTCt CAATAATGCT CCTGGACAGT 480
 44 ggCTCTGAGC agAACCGGCAg tGTCACATCA tGCTTAGAGC tGAATCTCTA tAAAATTGCT 540
 46 aAGCTGCAGA cCATGAACtA tATTGCTTG tGTTGTTGGCT GCCTGCTGCC ATTtTTcACA 600
 48 cTCAGCATCT GTTATCTGT GATCATTGCG GTTCTGTAA aAGTGGAGGT CCCAGAAATCG 660
 50 gggCTGCGGG tttCTCAGAG gAAGGCACTG ACCACCATCA tCATCACCTT GATCATCTTC 720
 52 ttCTCTGTGTT tCCTGCCCTA tCACACACTG AGGACCGTCC ACTTGACGAC atGGAAAGTG 780
 54 gGTTATGCA aAGACAGACT GcATAAAAGT ttGTTTATCA cACTGGCCTT GGCAGCAGCC 840
 56 aATGCTGTCT tCAATCCTCT GCTCTATTAC ttGCTGGGG AGAATTtAA ggACAGACTA 900
 58 aAGTCTGcac tCAGAAAAGG cCATCCACAG aAGGCAAAGA cAAAGTGTGT tttCCCTGTT 960
 60 agtGTGTGTT tgAGAAAGGA aACAAGAGTA tAA 993
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 65 <212> TYPE: PRT
 66 <213> ORGANISM: Homo sapiens
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 73 20 25 30
 75 Phe Phe Trp Gly Val Leu Gly Asn Gly Leu Ser Ile Tyr Val Phe Leu
 76 35 40 45
 78 Gln Pro Tyr Lys Lys Ser Thr Ser Val Asn Val Phe Met Leu Asn Leu
 79 50 55 60
 81 Ala Ile Ser Asp Leu Leu Phe Ile Ser Thr Leu Pro Phe Arg Ala Asp

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87	Ile	Met	Ser	Tyr	Ser	Leu	Tyr	Val	Asn	Met	Tyr	Ser	Ser	Ile	Tyr	Phe
88		100				105								110		
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91				115			120					125				
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94		130			135				140							
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97	145				150		155								160	
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105	Gly	Cys	Leu	Leu	Pro	Phe	Phe	Thr	Leu	Ser	Ile	Cys	Tyr	Leu	Leu	Ile
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112	225				230					235					240	
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124		290			295					300						
126	Arg	Lys	Gly	His	Pro	Gln	Lys	Ala	Lys	Thr	Lys	Cys	Val	Phe	Pro	Val
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163 1 5 10 15
165 atg gaa cca aat ggc acc ttc agc aat aac aac agc agg aac tgc aca 96
166 Met Glu Pro Asn Gly Thr Phe Ser Asn Asn Asn Ser Arg Asn Cys Thr
167 20 25 30
169 att gaa aac ttc aag aga gaa ttt ttc cca att gta tat ctg ata ata 144
170 Ile Glu Asn Phe Lys Arg Glu Phe Phe Pro Ile Val Tyr Leu Ile Ile
171 35 40 45
173 ttt ttc tgg gga gtc ttg gga aat ggg ttg tcc ata tat gtt ttc ctg 192
174 Phe Phe Trp Gly Val Leu Gly Asn Gly Leu Ser Ile Tyr Val Phe Leu
175 50 55 60
177 cag cct tat aag aag tcc aca tct gtg aac gtt ttc atg cta aat ctg 240
178 Gln Pro Tyr Lys Lys Ser Thr Ser Val Asn Val Phe Met Leu Asn Leu
179 65 70 75 80
181 gcc att tca gat ctc ctg ttc ata agc acg ctt ccc ttc agg gct gac 288
182 Ala Ile Ser Asp Leu Leu Phe Ile Ser Thr Leu Pro Phe Arg Ala Asp
183 85 90 95
185 tat tat ctt aga ggc tcc aat tgg ata ttt gga gac ctg gcc tgc agg 336
186 Tyr Tyr Leu Arg Gly Ser Asn Trp Ile Phe Gly Asp Leu Ala Cys Arg
187 100 105 110
189 att atg tct tat tcc ttg tat gtc aac atg tac agc agt att tat ttc 384
190 Ile Met Ser Tyr Ser Leu Tyr Val Asn Met Tyr Ser Ser Ile Tyr Phe
191 115 120 125
193 ctg acc gtg ctg agt gtt gtg cgt ttc ctg gca atg gtt cac ccc ttt 432
194 Leu Thr Val Leu Ser Val Val Arg Phe Leu Ala Met Val His Pro Phe
195 130 135 140
197 cgg ctt ctg cat gtc acc agc atc agg agt gcc tgg atc ctc tgt ggg 480
198 Arg Leu Leu His Val Thr Ser Ile Arg Ser Ala Trp Ile Leu Cys Gly
199 145 150 155 160
201 atc ata tgg atc ctt atc atg gct tcc tca ata atg ctc ctg gac agt 528
202 Ile Ile Trp Ile Leu Ile Met Ala Ser Ser Ile Met Leu Leu Asp Ser
203 165 170 175
205 ggc tct gag cag aac ggc agt gtc aca tca tgc tta gag ctg aat ctc 576
206 Gly Ser Glu Gln Asn Gly Ser Val Thr Ser Cys Leu Glu Leu Asn Leu
207 180 185 190
209 tat aaa att gct aag ctg cag acc atg aac tat att gcc ttg gtg gtg 624
210 Tyr Lys Ile Ala Lys Leu Gln Thr Met Asn Tyr Ile Ala Leu Val Val
211 195 200 205
213 ggc tgc ctg ctg cca ttt ttc aca ctc agc atc tgt tat ctg ctg atc 672
214 Gly Cys Leu Leu Pro Phe Phe Thr Leu Ser Ile Cys Tyr Leu Leu Ile
215 210 215 220
217 att cgg gtt ctg tta aaa gtg gag gtc cca gaa tcg ggg ctg cgg gtt 720
218 Ile Arg Val Leu Leu Lys Val Glu Val Pro Glu Ser Gly Leu Arg Val
219 225 230 235 240

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230	Thr	Trp	Lys	Val	Gly	Leu	Cys	Lys	Asp	Arg	Leu	His	Lys	Ala	Leu	Val		
231					275				280					285				
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268	Gln	Pro	Tyr	Lys	Lys	Ser	Thr	Ser	Val	Asn	Val	Phe	Met	Leu	Asn	Leu		
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292 Tyr Lys Ile Ala Lys Leu Gln Thr Met Asn Tyr Ile Ala Leu Val Val
293 195 200 205
295 Gly Cys Leu Leu Pro Phe Phe Thr Leu Ser Ile Cys Tyr Leu Leu Ile
296 210 215 220
298 Ile Arg Val Leu Leu Lys Val Glu Val Pro Glu Ser Gly Leu Arg Val
299 225 230 235 240
301 Ser His Arg Lys Ala Leu Thr Thr Ile Ile Ile Thr Leu Ile Ile Phe
302 245 250 255
304 Phe Leu Cys Phe Leu Pro Tyr His Thr Leu Arg Thr Val His Leu Thr
305 260 265 270
307 Thr Trp Lys Val Gly Leu Cys Lys Asp Arg Leu His Lys Ala Leu Val
308 275 280 285
310 Ile Thr Leu Ala Leu Ala Ala Asn Ala Cys Phe Asn Pro Leu Leu
311 290 295 300
313 Tyr Tyr Phe Ala Gly Glu Asn Phe Lys Asp Arg Leu Lys Ser Ala Leu
314 305 310 315 320
316 Arg Lys Gly His Pro Gln Lys Ala Lys Thr Lys Cys Val Phe Pro Val
317 325 330 335
319 Ser Val Trp Leu Arg Lys Glu Thr Arg Val
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VERIFICATION SUMMARY

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Input Set : A:\Pc10914a.app

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L:9 M:270 C: Current Application Number differs, Replaced Application Number

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date